APPROVED (O.C. F. 3.	CLASS SULCLASS	Which are an annual control of the c
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Analysis of 17724 (399 aa)

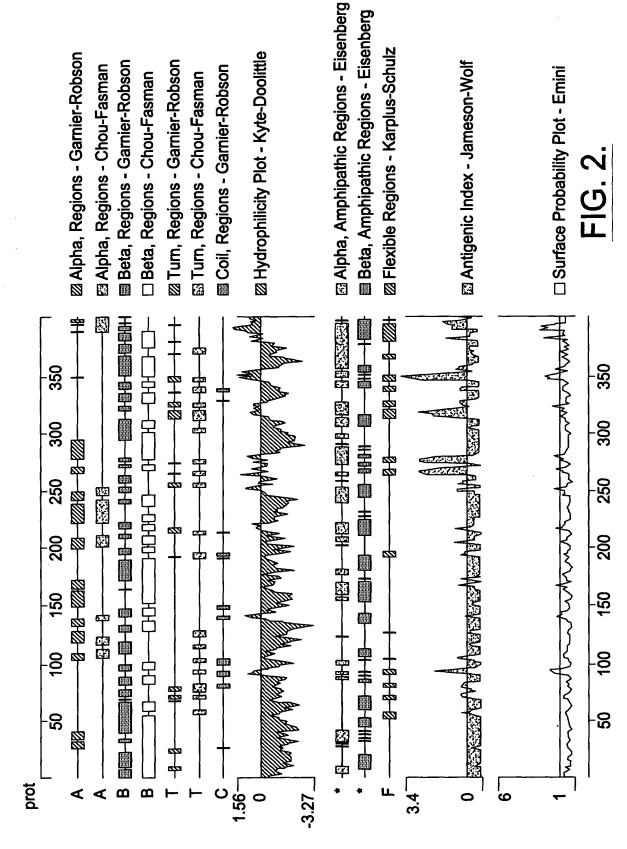
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FIG. 1.

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN Inventor(s): Glucksmann et al.

Application No: 09/78 1(380). | [... [...] |]... COUPLED RECEPTORS

Atty Dkt No: 35800/208932 (5800-206)



Atty Dkt No: 35800/208932 (5800-206)

3/22

Query: 17724 Model

Scores for sequence family classification (score includes all domains):

Description 7tm 1 7 transmembrane receptor (rhodopsin family)

Score E-value N 94. 1 6. 1e-29

Parsed for domains:

Model hmm-f hmm-t Domain seq-f seq-t score E-value 374 ... 259 () 94. 1 6. 1e-29 7tm_1 125 1

Alignments of top-scoring domains 7tm_1 : domain 1 of 1, from 125 to 374: score 94.1, E = 6.1e-29

> *->GN(LV: lv: lrtkk(rtptn:fi(NLAvADLLf(ltlppwalyylvg

GNTIIIVMVIADTHLHTPMYFFLGNFSLLEIŪVTMTAVPRMLSDLLV 171 17724 125

gsedWpfGsalCklvtaldvvnmyaSillLtaISiDRYlAIvhPlryrrr ++++ +C ++ ++ + ++ S | Lt +++DR++AI+hPlry ++

172 -- PHKVITFTGCMVQFYFHFSLGSTSFLILTDMALDRFVAICHPLRYGTL 219 17724

rtsprrAkvvillvWvlalllslPpllfswvktveegngtlnvnvtvCli

17724

17724

...kaaktllvvvvvFvlCWlPyfivllldtlc.lsiimsstCelervlp ++ + a+ ++ +++ v+ + i+l+++ + s ++ 316 sscQKAFSTCGSHLTLVFIGYSSTIFLYVRPGKaHS------VQ

17724

tallvtlwLayvNsclNPiIY<-*

+ v + (+ + + + + (NP + I)354 VRKVVALVTSVLTPFLNPFIL 374 17724

FIG. 3.

SOMMON

APPAC: .

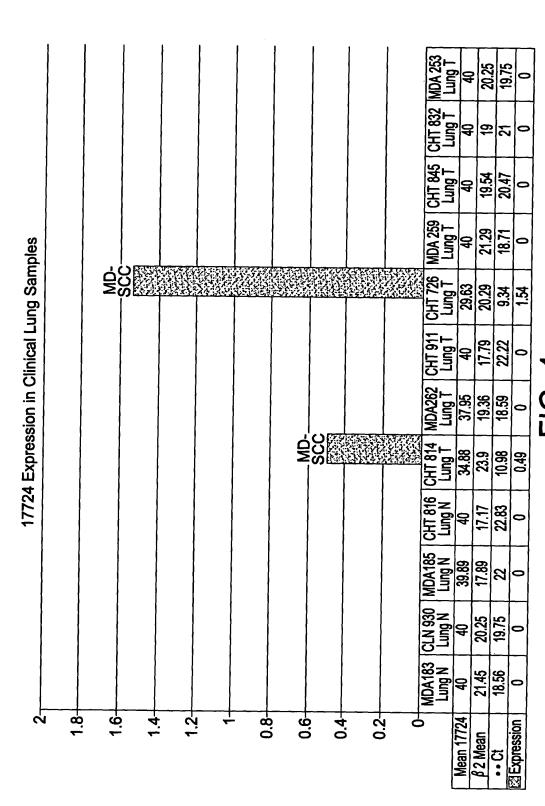


FIG. 4

SULLASS

ļ						Fetal Liver	39.8	20.1	8	
:							0.04	22.0	88	
						HMVEC-Arr HMVEC-Prol Fetal Liver	40.0	50.9	00	
						HMVECAL	40.0	18.5	80	
nples						Brain T	40.0	22.7	89	
17724 Expression in Clinical Anglogenic Samples						Brain T	40.0	21.6	8	
Angloge						Brain T	39.1	21.1	89	L
Clinical	!					Brain T	32.2	16.8	8	
sion in							38.4	21.2	83	
Expres	TORSE VAN OUT			2011	and and a wider	Astrocytes	38.5	50.9	8	
17724							30.9	23.2	4.8	
						Brain	32.2	24.8	00	
						Brain N	32.5	23.0	1.4	:
!						Brain N	40.0	23.6	8	İ L
ú		4. c	ა. -	, 4 - -	<u> </u>	0.0	17724	Beta 2	E-Expression	l İ

-1G. 5

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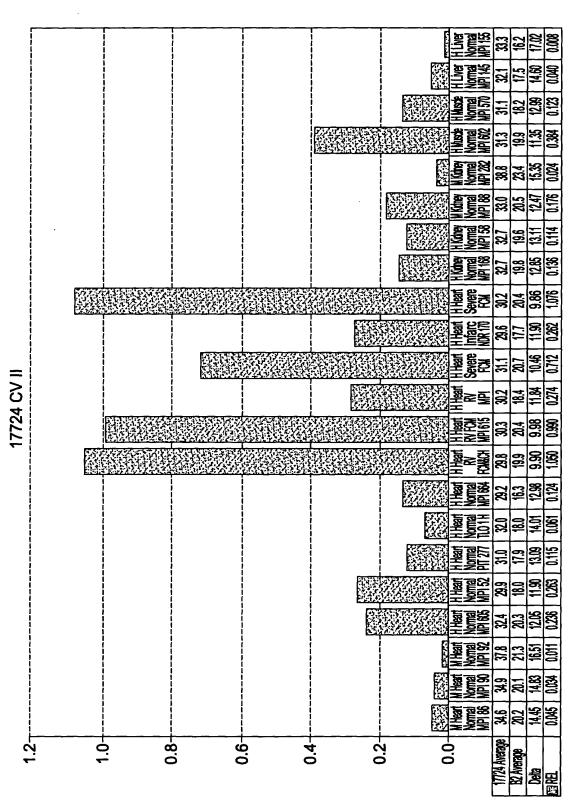


FIG. 6.

APPRIC D

BY

F
H Aorta H Aorta M M Normal Normal Aorta Aorta
8.4 4.4
20.0 19.2
0.046 0.230

FIG. 7

TO FIG. 8B. SJUCLASS 18.39 17.76 19.31 18.40 18.22 20.91 20.13 18.45 20.20 17.15 18.45 17.33 19.71 18.35 17.03 18.20 17.31 17.55 19.25 17.55 19.59 18.25 18.59 18.57 20.25 18.59 17.20 17.25 17.59 19.54 18.39 19.31 18.32 18.35 17.55 19.39 17.55 20.39 18.37 18.47 18.45 18.39 19.55 15.50 18.34 15.50 18.37 19.34 19.51 18.34 17.55 15.74 FIG. 8A. APLOND D.C. DHAF ST. 苔 <u>17.27 | 18.29 | 18.39 | 30.66 | 24.76 | 31.69 | 24.76 | 32.66 | 34.86 | 35.23 | 24.86 | 38.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05 | 36.05</u> Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN Phase 1.2.1 expression of 17724 8 COUPLED RECEPTORS 21.96 18.99 17.76 19.31 18.40 18.23 20.91 20.13 18.45 20.20 1 37.76 31.01 33.16 33.84 34.20 35.70 Aorta/Fetal | norma heart 0.5 3.0-2.0-<u>P</u>. 2.5 1.5-E-Emeson

ВY Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN COUPLED RECEPTORS

FROM FIG. 8A.

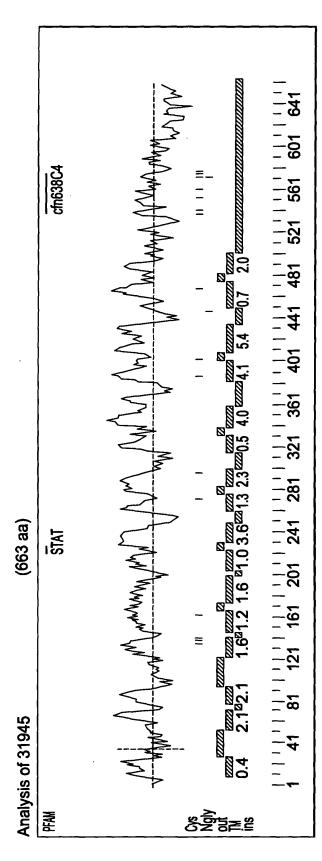
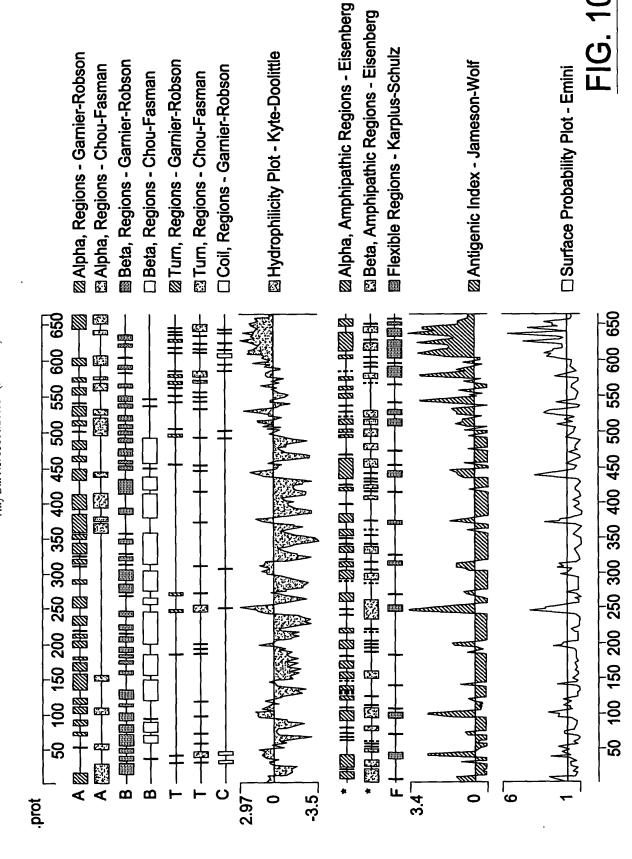


FIG. 9.

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN Inventor(s): Glucksmann et al. COUPLED RECEPTORS



APPLICATED 1 O.C.

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN

à

Inventor(s): Glucksmann et al.

COUPLED RECEPTORS

23.2 29.3 22.4 30.8 22.0 20.6 19.8 20.0 20.4 24.8 21.6 40. 24.8 28.3 29.4 30.2 26.6 26.5 26.8 27.5 28.1 30.1 30.0 32.9 27.5 32.8 26.9 25.5 25.3 24.9 25.5 28.1 90.2 85.9 57.3 91.1 107.3 291.0 20.9 45.7 92.4 112.2 173.0 301.3 500.0 33.6 23.9 222 Taqman Chart+Table GPCR 31945 Expression 20.1 25.1 22.4 24.4 21.8 21.9 22.9 24.3 25.9 23.7 Tă M NHLF hango MAKES.) Mg **W** W. 8 **88**8 112.6 69.8 166.0 20.5 25.3 21. 25.3|26.4|29.2|25.1|30.6|25. 636.9 528.2 482.7 33) SE SE Relative Expression (TH2 24 RLD63 used as reference sampl) 600.0− 500.0-400.0-300.0-100.0-E Egression 1500.0-1400.0-¥g

FIG. 11A.

A PATOMED 1.0

NETTS: N

(会) 25.4 26.2 25.3 24.7 24.8 25.5 24.6 24.9 31.0 29.3 26.9 29.6 25.8 26.9 25.4 26.9 29.7 29.9 26.0 18.5 20.4 18.4 18.9 18.9 24.6 23.6 23.3 19.6 25.4 25.4 23.6 21.8 24.4 22.4 23.9 23.8 21.7 21.1 22.5 48.7 23.2 48.8 43.4 1423 1331 331.8 31.0 176.1 955.4 44.6 170.7 477.7 327.4 336.6 43.9 89 92.1 開き 8 害 PBMC NHBE NHBE 24hr (RLD6 BM- mPB ABM mBM TNP, mock 1113-1 (L67) 3) MNC 0334+ 0334+ 0334+

FROM FIG. 11A.

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN COUPLED RECEPTORS

Inven Glucksmann et al. Appli No: 09/781,880

Atty Dkt No: 35800/208932 (5800-206)

14/22

Analysis of 50288 (372 aa)

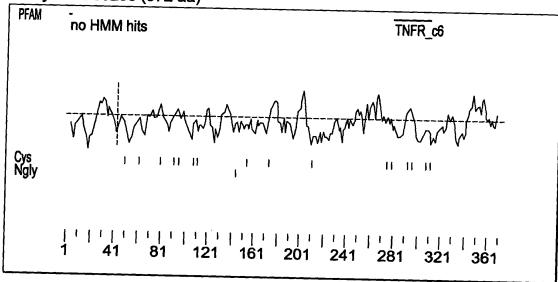


FIG. 12.

DSFELETO . Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN Application No: 09/781 880 mm [mm] mm Atty Dkt No: 35800/208932 (5800-206) Inventor(s): Glucksmann et al. COUPLED RECEPTORS

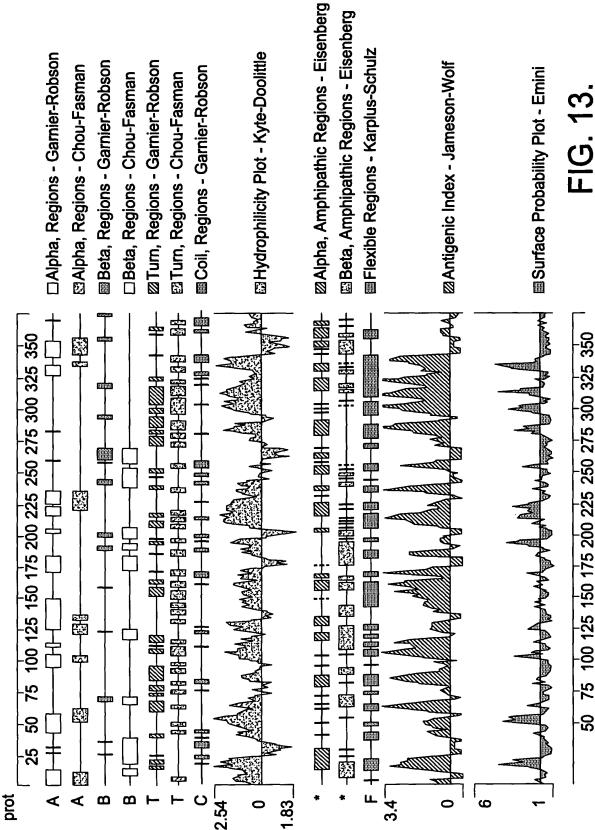


FIG. 13

CENCUED.

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN

Inventor(s): Glucksmann et al.
Application No. 09/781,880 [] T. T. [] T.

COUPLED RECEPTORS

Atty Dkt No: 35800/208932 (5800-206)

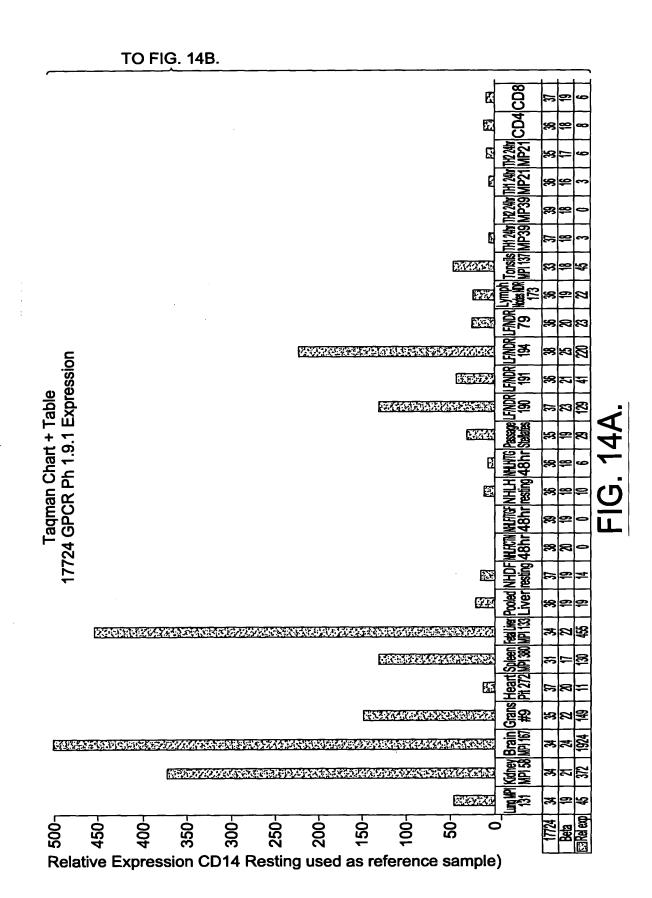


FIG. 14B.

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E	mBM CORRECT 16-	æ	æ	∞
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	4 PBM g Moc	æ	┢	4
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FROM FIG. 14A.

TO FIG. 15B.

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											NDR NDR 07 NDR 12 PIT 208 CHT 620 CHT 619 MA CLN 03 CLN 05 CLN 17 CLN 07 CLN 08 MA CLN 02 MDA 25 Breast T Breast T Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ovary N Ova	9	\$	2	9	
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ressi											25.58 1.58 1.58 1.58 1.59 1.59	3.13	213	38 88	0	
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7724											Pesset T	8	9	ä	0	
											\$ 55 kg	37.98	40	\$	0	
										R 22	\$\$\$ \$\$	386		83	8	
											PIT 56 Breast N	8	125	\$	0	
											PTT 371 Breast N	40	23.95	3	0	
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0.5	0.45	0.4	0.35	0.3	0.25	0.2-	0.15	0.1	0.05	C	5	Hean 17724	821831	3 3 3	ESEQUESSION	

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN COUPLED RECEPTORS Inventor(s): Glucksmann et al. Application No: 09/781,880 []] T. T. []] T. " []] []] [] (]] Atty Dkt No: 35800/208932 (5800-206)

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN COUPLED RECEPTORS Inventor(s): Glucksmann et al. Application No. 09[[2]] [2] [2] [2] [2] [2] [2] [2] [3] [4] [4] Atty Dkt No: 35800/208932 (5800-206)

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	重氮		8	77.30	1871	0	
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	CHF911		40	(I.N	22	0	5B
	€ §	Ę	37.55	<u> </u>	63	0	
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	₹		3	3	\$	0	
· · · · · · · · · · · · · · · · · · ·			Н	-	-	\vdash	

FROM FIG. 15A.

TO FIG. 16B.

MPI 613 | NDR 170 | MPI 617 | MPI 168 | MPI 58 | MPI 88 | MPI 282 | MPI 602 | MPI 570 | MPI 145 | MPI 155 | $\left| 0.034 \left| 0.011 \right| 0.236 \right| 0.263 \right| 0.115 \right| 0.061 \right| 0.124 \right| 1.050 \right| 0.990 \right| 0.274 \right| 0.712 \right| 0.262 \right| 1.076 \right| 0.136 \right| 0.114 \right| 0.176 \right| 0.024 \right| 0.384 \right| 0.123 \right| 0.040 \right| 0.008 \right| 0.300 \right| 0.034 \right| 0.123 \right| 0.040 \right| 0.008 \right| 0.300 \right| 0.034 \\ 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \\| 0.036 \right| 0.036 \right| 0.036 \right| 0.036 \\| 0.036 \right$ FIG. 16A 14.45 | 14.83 | 16.51 | 12.05 | 11.90 | 13.09 | 14.01 | 12.98 | 9.90 | 9.98 | 11.84 | 10.46 | 11.90 | 9.86 | 12.85 | 13.11 | 12.47 | 15.35 | 11.35 | 12.99 | 14.45 | 14.81 | 12.47 | 15.35 | 14.35 | 12.99 | 14.45 | 14.81 | 12.47 | 15.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 | 14.35 |38.8 33.0 32.7 32.7 30.2 29.6 17724 CV II and III 31.1 Inventor(s): Glucksmann et al.
Application No: 09/78 [[380]]. | | | | | | | | | | | | | | | Atty Dkt No: 35800/208932 (5800-206) 30.2 MPI 614 MPI 615 30.3 29.8 MPI 86 | MPI 90 | MPI 92 | MPI 605 | MPI 52 | PIT 277 | 71.0 1 H | MPI 664 | 29.2 32.0 34.9 34.6 20.2 0 4-3-4 5 9 图0.045

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN

配 355 26.6 11.38 12.70 8.13 12.45 10.31 4.98 9.54 10.94 7.95 8 9 9 **医** 31.2 N.W. 怒 28.7 3 31 33.2 MP1 526 | MP1 442 | MP1 438 | MP1 532 | CLN 67 | CLN 66 | 40.0 0 34.2 26.1 33.5 33.6 33.1 MPI 525 34.3 25.0 0 MPI 93 MPI 528 31.3 34.4 4.4 12.42 14.41 CLN 618 PTT 394 35.6 35.0

FROM FIG. 16A.

Title: NOVEL SEVEN-TRANSMEMBRANE PROTEIN/G-PROTEIN Application No. 09/781,880 Atty Dkt No. 35800/208932 (5800-206) Inventor(s): Glucksmann et al. COUPLED RECEPTORS

